CERTIFICATE OF MAILING 37 C.F.R. 1.8

I her by certify that this correspond ince is biling deposited with the U.S. Postal Service with sufficient postage as First Class Mail in an envelope addressed to: Commission r for Patents, Washington, DC 20231, on th date b low:



PATENT

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of:

Lieven Stuvver Joost Louwagie

Rudi Rossau

Serial No.: 09/943.983

Filed: August 31, 2001

For: METHOD FOR DETECTION OF DRUG-

INDUCED MUTATIONS IN THE REVERSE TRANSCRIPTASE GENE

firmation No.:

oup Art Unit: 1656

Examiner: To be assigned

Atty. Dkt. No.: 11362.0008.DVES02

JAN 22

Live

T (b) 100 STATEMENT AS REQUIRED UNDER 37 C.F.R. § 1.825(a) AND (b) AND STATEMENT AS REQUIRED UNDER 37 C.F.R. § 1.821(g)

BOX SEQUENCE

Commissioner for Patents Washington, D.C. 20231

Sir:

Submitted herewith is a substitute computer readable form and a substitute paper copy of the sequence listing of those sequences in the captioned patent application, wherein minor errors have been corrected. The substitute computer readable form of the sequence listing is the same as the substitute paper copy of the sequence listing. The sequence information provided in the Specification is also the same as the sequence listing of the enclosed substitute computer Please direct all future communications regarding this application as follows:

Matthew L. Madsen HOWREY SIMON ARNOLD & WHITE, LLP 750 Bering Drive Houston, Texas 77057-2198 (713) 787-1400

Respectfully submitted,

Patricia A. Kammerer

Reg. No. 29,775 Attorney for Assignee INNOGENETICS N.V.

HOWREY SIMON ARNOLD & WHITE, LLP 750 Bering Drive Houston, Texas 77057-2198 (713) 787-1400

Dav. 5, 2001



Patentin 2.0

"bue"

Misuse of n

Raw Sequence Listing Error Summary

ORIGINAL INTERES

SERIAL NUMBER: 09/943,983 ERROR DETECTED SUGGESTED CORRECTION ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPIIA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE Wrapped Nucleics The number/text at the end of each line "wrapped" down to the next line. This may occur if your file Wrapped Aminos was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping." The rules require that a line not exceed 72 characters in length. This includes white spaces. Invalid Line Length Misaligned Amino The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; Numbering use space characters, instead. Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCI1 text. Sequence(s)___contain n's or Xaa's representing more than one residue. Per Sequence Rules, Variable Length. each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing. Patentin 2.0 A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid "bug" . Normally, Patentin would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences. Skipped Sequences missing. If intentional please insert the following lines for each skipped sequence: (OLD RULES) (2) INFORMATION FOR SEO ID NO:X: (insert SEO ID NO where "X" is shown) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to Include the skipped sequences. missing. If Intentional, please insert the following lines for each skipped sequence. Sequence(s) ___ Skipped Sequences (NEW RULES) <210> sequence id number <400> sequence id number 000 Use of n's or Xaa's Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. (NEW RULES) In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents. Invalid <213> Per 1,823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or Response scientific'name (Genus/species). <220>-<223> section is regulred when <213> response is Unknown or scienture name versions is Artificial Sequence (A mode) Feature" and associated numeric identifiers and responses. Security () - 200 missing the 4200 "Feature" and associated numeric identifiers and responses. Use of <220> "Unknown." Please explain source of genetic material in <220> to <223> section.

in can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent AMC/MH - Biotechnology Systems Branch - 08/21/2001

any value not specifically a nucleotide.

(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)

Please do not use "Copy to Disk" function of Patentin version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence

listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.





OIPE

RAW SEQUENCE LISTING PATENT APPLICATION: US/09/943,983

DATE: 09/20/2001 TIME: 13:08:01

Input Set : A:\ES.txt

Output Set: N:\CRF3\09202001\I943983.raw

Does Not Comply

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Corrected Diskette Needed
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             Louwagie, Joost
             Rossau, Rudi
     7 <120> TITLE OF INVENTION: METHOD FOR DETECTION OF DRUG-INDUCED MUTATIONS IN THE
EVERSE
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    10 <130> FILE REFERENCE: 11362.0008.DVUS02 INNS008--3
 -> 12 <140> CURRENT APPLICATION NUMBER: US/09/943,983
12 (141) CURRENT FILING DATE: 2001-08-31
    15 <150> PRIOR APPLICATION NUMBER: 48/913,833 now US/6,087,093
    16 <151> PRIOR FILING DATE: 1997-09-15
    18 <150> PRIOR APPLICATION NUMBER: PCT/EP 97/00211
    19 <151> PRIOR FILING DATE: 1997-01-17
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    25 <151> PRIOR FILING DATE: 1996-06-25
    27 <160> NUMBER OF SEQ ID NOS: 164
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55 aaatggaaaa ggaag

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15

RAW SEQUENCE LISTING

DATE: 09/20/2001 TIME: 13:08:01 PATENT APPLICATION: US/09/943,983

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Output Set: N:\CRF3\09202001\I943983.raw

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--> 117 <220> FEATURE: --> 117 (223) OTHER INFORMATION:

115 ORGANISM Artificial sequence

RAW SEQUENCE LISTING

DATE: 09/20/2001 TIME: 13:08:01 PATENT APPLICATION: US/09/943,983

Input Set : A:\ES.txt

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Output Set: N:\CRF3\09202001\I943983.raw
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/943,983

DATE: 09/20/2001 TIME: 13:08:01

Input Set : A:\ES.txt

Output Set: N:\CRF3\09202001\I943983.raw

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226 acagaactgg aaaa

14

DATE: 09/20/2001 TIME: 13:08:01

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                      Output Set: N:\CRF3\09202001\1943983.raw
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/943,983

The types of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/943,983

DATE: 09/20/2001 TIME: 13:08:02

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- Output Set: N:\CRF3\09202001\1943983.raw :12 M:270 C: Current Application Number differs, Replaced Current Application Number :13 M:271 C: Current Filing Date differs, Replaced Current Filing Date ::36 M:258 W: Mandatory Feature missing, <220> FEATURE: :36 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: ::45 M:258 W: Mandatory Feature missing, <220> FEATURE: :45 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: ::54 M:258 W: Mandatory Feature missing, <220> FEATURE: ::54 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: ::63 M:258 W: Mandatory Feature missing, <220> FEATURE: ::63 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: ::72 M:258 W: Mandatory Feature missing, <220> FEATURE: :72 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: ::81 M:258 W: Mandatory Feature missing, <220> FEATURE: :81 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: .:90 M:258 W: Mandatory Feature missing, <220> FEATURE: :90 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: ::99 M:258 W: Mandatory Feature missing, <220> FEATURE: :99 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: ::108 M:258 W: Mandatory Feature missing, <220> FEATURE: ::108 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: ::117 M:258 W: Mandatory Feature missing, <220> FEATURE: ::117 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: ::126 M:258 W: Mandatory Feature missing, <220> FEATURE: :: 126 M: 258 W: Mandatory Feature missing, <223> OTHER INFORMATION: ::135 M:258 W: Mandatory Feature missing, <220> FEATURE: ::135 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: ::144 M:258 W: Mandatory Feature missing, <220> FEATURE: :144 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: ::153 M:258 W: Mandatory Feature missing, <220> FEATURE: :153 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: ::162 M:258 W: Mandatory Feature missing, <220> FEATURE:
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VERIFICATION SUMMARYDATE: 09/20/2001PATENT APPLICATION: US/09/943,983DATE: 09/20/2001TIME: 13:08:02

Input Set : A:\ES.txt

Output Set: N:\CRF3\09202001\1943983.raw

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